Rec'd PCT/PTO 09 SEP 2004



10/506549

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/506,549

DATE: 09/09/2004 TIME: 16:15:50

Input Set : A:\SEQLIST 1361US.txt

Output Set: N:\CRF4\09092004\J506549.raw

```
4 <110> APPLICANT: APPLERA CORPORATION
      6 <120> TITLE OF INVENTION: ISOLATED HUMAN TRANSPORTER PROTEINS,
             NUCLEIC ACID MOLECULES ENCODING HUMAN TRANSPORTER PROTEINS,
             AND USES THEREOF
     10 <130> FILE REFERENCE: CL001361-US
C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/506,549
C--> 13 <141> CURRENT FILING DATE: 2004-09-03
     15'<150> PRIOR APPLICATION NUMBER: 60/361,343
    16 <151> PRIOR FILING DATE: 2002-03-05
    18 <160> NUMBER OF SEQ ID NOS: 4
```

20 <170> SOFTWARE: FastSEQ for Windows Version 4.0 22 <210> SEQ ID NO: 1

23 <211> LENGTH: 2679 24 <212> TYPE: DNA

25 <213 > ORGANISM: Homo sapiens 27 <400> SEQUENCE: 1 28 atgtcttcca agaagaatag aaagcggttg aaccaaagcg cggaaaatgg ttcgtccttg 60 29 ccctctgctg cttcctcttg tgcggaggca cgggctcctt ctgctggatc agacttcgcg 120 30 gcaacctccg ggactctgac ggtgaccaac ttattagaaa aggatgacaa aattcctaaa 180 31 acattccaga attcccttat tcatcttgga ctcaacacta tgaagtctgc aaatatatgt 240 32 ataggtcgac cagtgttgct tactagtttg aacggaaagc aagaggtgta tacagcctgg 300 33 cctatggcag gatttcctgg aggcaaggtc ggcctgagtg aaatggcaca gaaaaatgtg 360 34 ggtgtgaggc ctggtgatgc catccaggtc cagcctcttg tgggtgctgt gctacaggct 420 35 gaggaaatgg atgtggcact gagtgacaaa gatatggaaa ttaatgaaga agaactgact 480 36 ggttgtatcc tgagaaaact agatggcaag attgttttac caggcaactt tctgtattgt 540 37 acattctatg gacgaccgta caagctgcaa gtattgcgag tgaaaggggc agatggcatg 600 38 atattgggag ggcctcagag tgactctgac actgatgccc aaagaatggc ctttgaacag 660 39 tocagcatgg aaaccagtag cotggagtta toottacago taagccagtt agatotggag 720 40 gatacccaga teccaacate aagaagtaet eettataaae caattgatga cagaattaca 780 41 aataaagcca gtgatgtttt gctggatgtt acacagagcc ctggagatgg cagtggactt 840 42 atgctagagg aagtcacagg tcttaaatgt aattttgaat ctgccagaga aggaaatgag 900 43 caacttactg aagaagagag actgctaaag ttcagcatag gagcaaagtg caatactgat 960 44 actttttatt ttatttcttc aacaacaaga gtcaatttta cagagattga taaaaattca 1020 45 aaaqaqcaaq acaaccaatt caaaqtaact tatqacatga tagqaqgatt aagtagccag 1080 46 ctgaaagcaa ttagagaaat aattgaattg cccctcaaac agcctgagct tttcaagagt 1140 47 tatggaattc ctgcccctag aggagtgtta ctttatggtc ctccaggtac tggaaaaaca 1200 48 atgatcgcca gggctgttgc taatgaagtt ggagcctatg tttctgtaat taatggtcct 1260 49 qaaattataa qcaaattcta tqqtqaqact qaaqcaaaqt tacqtcaqat atttqctgaa 1320 50 gccactctac gacacccatc aattattttt attgatgagc tggatgcact ttgtccgaaa 1380 51 agagaggggg cccagaatga agtggaaaaa agagttgtgg cttcactctt aacactgatg 1440 52 gatggcattg gttcagaagt aagtgaagga caagtgttgg ttcttggggc cacaaatcgc 1500 53 cctcatgcct tggatgctgc tctccgaaga cctgggcgat ttgataaaga gattgagatt 1560

54 ggagttccca atgctcagga ccggctagat attctccaga aactgcttcg aagggtaccc 1620

Input Set : A:\SEQLIST_1361US.txt
Output Set: N:\CRF4\09092004\J506549.raw

```
55 catttgctca ctgaggctga gctgctgcag ctggcaaata gtgctcatgg atacgttgga 1680
56 gcagacttga aagtcttgtg taatgaagca ggtctctgtg ccttgcggag aatcctgaaa 1740
57 aaacagccta acctccctga tgtcaaggtg gctggactgg tgaagattac tctgaaggat 1800
58 ttcttgcagg caatgaatga tatcagaccc agtgccatga gggaaatagc aattgatgtc 1860
59 ccaaatgtat cctggtcaga tataggagga ctggaaagta tcaaactgaa gttggaacag 1920
60 gctgtggaat ggcccttaaa acatccagag tctttcattc gaatgggtat tcagccacct 1980
61 aaaggagttc ttctctatgg gccacctggg tgctctaaaa caatgatagc aaaggctttg 2040
62 gccaatgaga gtggactgaa ttttctagct ataaaggggc ctgaattaat gaataaatat 2100
63 gttggtgaat ctgaaagagc agttagagag accttccgaa aagcaagagc agtggcgcct 2160
64 tocattattt totttgatga actggatgco ttagcagttg aaaggggcag ttotttaggt 2220
65 qctqqqaatq taqccqatcq tqttttqqct caqctcttaa caqaaatqqa tqqqattqaa 2280
66 cagctaaagg atgtgaccat tttggcagct actaaccgtc cagataggat agacaaggct 2340
67 ttgatgcggc ctggaagaat tgatagaatc atctatgtgc ctttaccgga tgcagcaaca 2400-
68 agaagggaaa tatttaagct gcagtttcac tccatgcctg tcagtaatga agttgacctg 2460
69 gatgaactca teetteaaac egaegeatae teaggageag agattgtage tgtetgeaga 2520
70 gaggcagctc ttctggctct ggaagaagac attcaagcca atctcatcat gaaaagacat 2580
71 ttcactcagg ccttgagcac tgtgacacct agaattcctg agtcattgag acgtttttat 2640
72 gaagattatc aagagaagag tgggctgcat acactctga
                                                                      2679
74 <210> SEQ ID NO: 2
75 <211> LENGTH: 892
76 <212> TYPE: PRT
77 <213> ORGANISM: Homo sapiens
79 <400> SEQUENCE: 2
80 Met Ser Ser Lys Lys Asn Arg Lys Arg Leu Asn Gln Ser Ala Glu Asn
                                       10
82 Gly Ser Ser Leu Pro Ser Ala Ala Ser Ser Cys Ala Glu Ala Arq Ala
               20
                                   25
84 Pro Ser Ala Gly Ser Asp Phe Ala Ala Thr Ser Gly Thr Leu Thr Val
85
86 Thr Asn Leu Leu Glu Lys Asp Asp Lys Ile Pro Lys Thr Phe Gln Asn
87
                           55
88 Ser Leu Ile His Leu Gly Leu Asn Thr Met Lys Ser Ala Asn Ile Cys
                       70
                                           75
90 Ile Gly Arg Pro Val Leu Leu Thr Ser Leu Asn Gly Lys Gln Glu Val
92 Tyr Thr Ala Trp Pro Met Ala Gly Phe Pro Gly Gly Lys Val Gly Leu
               100
                                   105
94 Ser Glu Met Ala Gln Lys Asn Val Gly Val Arg Pro Gly Asp Ala Ile
                               120
                                                   125
          115
96 Gln Val Gln Pro Leu Val Gly Ala Val Leu Gln Ala Glu Glu Met Asp
                           135
98 Val Ala Leu Ser Asp Lys Asp Met Glu Ile Asn Glu Glu Glu Leu Thr
                       150
100 Gly Cys Ile Leu Arg Lys Leu Asp Gly Lys Ile Val Leu Pro Gly Asn
101
                                        170
                    165
102 Phe Leu Tyr Cys Thr Phe Tyr Gly Arg Pro Tyr Lys Leu Gln Val Leu
                180
                                    185
104 Arg Val Lys Gly Ala Asp Gly Met Ile Leu Gly Gly Pro Gln Ser Asp
                                200
105
            195
                                                    205
```

Input Set : A:\SEQLIST_1361US.txt

Output Set: N:\CRF4\09092004\J506549.raw

106 107		Asp 210	Thr	Asp	Ala	Gln	Arg 215	Met	Ala	Phe	Glu	Gln 220	Ser	Ser	Met	Glu
108	Thr	Ser	Ser	Leu	Glu	Len	Ser	Len	Gln	Leu	Ser	Gln	Len	Asp	Len	Glu
	225					230					235	01		1101		240
		Thr	Cln	т10	Dro		Cox	7~~	502	The		Ф	T	Dwo	T1.	
111	Asp	1111	GIII	iie	245	IIII	Ser	Arg	Ser	250	PIO	īĀī	гуѕ	Pro	255	Asp
112	Asp	Arg	Ile	Thr	Asn	Lys	Ala	Ser	Asp	Val	Leu	Leu	Asp	Val	Thr	Gln
113				260					265				-	270		
114	Ser	Pro	Glv	Asp	Glv	Ser	Gly	Leu	Met	Leu	Glu	Glu	Val	Thr	Glv	Leu
115			275	_			2	280					285		1	
	Lvs	Cvs		Phe	Glu	Ser	Ala		Glu	Glv	Δen	Glu		T.011	Thr	Glu
117	27.5	290	11011	1110	014	UCI	295	9	OLU	GLY	ASII	300	GIII	Leu	1111	GIU
	C1		7	T	T	T		C	T1.	a 1	71-		C	7	m1	3
		Gru	Arg	ьeu	ьeu	_	Phe	Ser	тте	GIY		гуѕ	Cys	Asn	Tnr	_
	305		_		_	310		_	_		315				_	320
	Thr	Phe	Tyr	Phe		Ser	Ser	Thr	Thr	_	Val	Asn	Phe	Thr	Glu	Ile
121					325					330					335	
122	Asp	Lys	Asn	Ser	Lys	Glu	Gln	Asp	Asn	Gln	Phe	Lys	Val	\mathtt{Thr}	Tyr	Asp
123				340					345					350		
124	Met	Ile	Gly	Gly	Leu	Ser	Ser	Gln	Leu	Lys	Ala	Ile	Arg	Glu	Ile	Ile
125			355					360		_			365			
	Glu	Leu	Pro	Leu	Lvs	Gln	Pro	Glu	Leu	Phe	Lvs	Ser	Tvr	Glv	Ile	Pro
127		370			•		375				*	380	•			
128	Ala	Pro	Ara	Glv	Val	Leu	Leu	Tvr	Glv	Pro	Pro		Thr	Glv	Lvs	Thr
	385		5			390		- 4 -	2		395	1		1	-1-	400
		Tle	Ala	Ara	Ala		Ala	Asn	G111	Val	Glv	Δla	Tvr	Val	Ser	
131				5	405				014	410	011		-1-	, 41	415	
	Tle	Δeń	Glv	Pro		Tla	Ile	Ser	Lvc		Туг	G137	Glu	Thr		λl = ·
133	110	ASII	Gry	420	GIU	110	ııç	Ser	425	FIIC	TYL	Gry	GIU	430	GIU	AIG
	Taro	T 011	7~~		т1.	Dho	Ala	C1	_	mb se	T 011	7~~	774 ~		C	T1.
	пур	ьеu	_	GIII	116	Pile	нта		Ата	TIIL	ьeu	Arg		PIO	ser	iie
135	T7 -	Dh -	435	3	~ 1	•	3	440	.	a .	.	_	445	~ 3	~3	
	тте		тте	Asp	GIU	Leu	Asp	Ата	ьeu	Cys	Pro	_	Arg	GIU	GIY	Ата
137	~ 1.	450	~7		~3	_	455				_	460	_	_,	_	
		Asn	GIU	vал	GIu	_	Arg	Val	Val	Ala		Leu	Leu	Thr	Leu	
	465					470	_			_	475	_		_		480
140	Asp	Gly	Ile	Gly	Ser	Glu	Val	Ser	Glu	_	Gln	Val	Leu	Val	Leu	Gly
141					485					490					495	
142	Ala	Thr	Asn	Arg	Pro	His	Ala	Leu	Asp	Ala	Ala	Leu	Arg	Arg	Pro	Gly
143				500					505					510		
144	Arg	Phe	Asp	Lys	Glu	Ile	Glu	Ile	Gly	Val	Pro	Asn	Ala	Gln	Asp	Arg
145			515					520					525			
146	Leu	Asp	Ile	Leu	Gln	Lys	Leu	Leu	Arg	Arg	Val	Pro	His	Leu	Leu	Thr
147		530				_	535		_	_		540				
148	Glu	Ala	Glu	Leu	Leu	Gln	Leu	Ala	Asn	Ser	Ala	His	Glv	Tvr	Val	Glv
149						550					555		3	- 4 -		560
		Asp	Leu	Lvs	Val		Cys	Asn	G] 11	Ala		Len	Cvs	Ala	Len	
151		<u>-</u> -		-1-	565		-1-			570	1		J, 5		575	- 5
	Ara	Tle	T.e.ii	Lvc		Gln	Pro	Δen	T.e.ii		Δen	₩-1	Lve	1721		Glv
153			 _u	580	-75	O-11	110	-1011	585	110	Ph	VAI	ny s	590	лта	O-Y
	Leu	v-1	Luc		ሞኮ∽	Len	Larg	7.00		Lev	Gl-	71-	Mot		7 c ~	т1 ^
104	TEU	val	пÄр	TTG	TIIT	neu	Lys	veb	FIIE	neu	GIII	нтq	MEL	ASII	Asp	тте

Input Set : A:\SEQLIST 1361US.txt

Output Set: N:\CRF4\09092004\J506549.raw

```
155
            595
                                600
156 Arg Pro Ser Ala Met Arg Glu Ile Ala Ile Asp Val Pro Asn Val Ser
        610
                            615
                                                 620
158 Trp Ser Asp Ile Gly Gly Leu Glu Ser Ile Lys Leu Lys Leu Glu Gln
                        630
                                            635
160 Ala Val Glu Trp Pro Leu Lys His Pro Glu Ser Phe Ile Arg Met Gly
                    645
                                        650
162 Ile Gln Pro Pro Lys Gly Val Leu Leu Tyr Gly Pro Pro Gly Cys Ser
                660
                                    665
164 Lys Thr Met Ile Ala Lys Ala Leu Ala Asn Glu Ser Gly Leu Asn Phe
            675
                    .
                                680
                                                     685
166 Leu Ala Ile Lys Gly Pro Glu Leu Met Asn Lys Tyr Val Gly Glu Ser
                           695
168 Glu Arg Ala Val Arg Glu Thr Phe Arg Lys Ala Arg Ala Val Ala Pro
                        710
                                             715
170 Ser Ile Ile Phe Phe Asp Glu Leu Asp Ala Leu Ala Val Glu Arg Gly
171
                                        730
                    725
172 Ser Ser Leu Gly Ala Gly Asn Val Ala Asp Arg Val Leu Ala Gln Leu
                                    745
174 Leu Thr Glu Met Asp Gly Ile Glu Gln Leu Lys Asp Val Thr Ile Leu
            755
                                760
176 Ala Ala Thr Asn Arg Pro Asp Arg Ile Asp Lys Ala Leu Met Arg Pro
                            775
178 Gly Arg Ile Asp Arg Ile Ile Tyr Val Pro Leu Pro Asp Ala Ala Thr
                                            795
179 785
                        790
180 Arg Arg Glu Ile Phe Lys Leu Gln Phe His Ser Met Pro Val Ser Asn
                    805
                                        810
182 Glu Val Asp Leu Asp Glu Leu Ile Leu Gln Thr Asp Ala Tyr Ser Gly
                820
                                    825
183
184 Ala Glu Ile Val Ala Val Cys Arg Glu Ala Ala Leu Leu Ala Leu Glu
185
                                840
186 Glu Asp Ile Gln Ala Asn Leu Ile Met Lys Arg His Phe Thr Gln Ala
                                                 860
187
                            855
188 Leu Ser Thr Val Thr Pro Arg Ile Pro Glu Ser Leu Arg Arg Phe Tyr
                        870
190 Glu Asp Tyr Gln Glu Lys Ser Gly Leu His Thr Leu
191
                    885
                                        890
194 <210> SEQ ID NO: 3
195 <211> LENGTH: 394191
196 <212> TYPE: DNA
197 <213> ORGANISM: Homo sapiens
199 <220> FEATURE:
200 <221> NAME/KEY: misc_feature
201 <222> LOCATION: (1)...(394191)
202 <223> OTHER INFORMATION: n = A, T, C or G
204 <400> SEQUENCE: 3
205 caagetttet tateattggg etcaaggttg aaagecagta aaaagtttte ttattgaaga 60
206 ctgaaaccac ccaccaccqt ccctqqttaa tcataaaaga agcgttattc taaaaaactc 120
207 caqttctcqc tqaqacaqct qttqacccaa tttgtataca aatqtgacac tctgaacctc 180
```

Input Set : A:\SEQLIST 1361US.txt

Output Set: N:\CRF4\09092004\J506549.raw

```
208 tctggtttag tatttgatag cccaacaggg tgactatatt caataaagat tcaggctggg 240
209 tgcagtggct cctgcctgta atcccagcac tttgaggggc caaggcggga gaatcagttg 300
210 agcccgggag actgagacca gcctgggcaa cacagtgaga caccgtttct agaaaaaata 360
211 gaaaaaagcc agatgcgcgc ctgtagtcct agctacccag gaggctgagg tgggaggatt 420
212 gcttgagctc aggggttgga ggctgcagtg aaccctgatc ctaccactgc actctagcct 480
214 taaaaataac taagagtaac tggattatgg ttgaatgctt gtggtgatga gtatctcatt 600
215 taccctgatg taattattac acatttatac ctctatcaaa atagctcata taccccataa 660
217 ccactcagtc caccacccgt attagaatgt aagagttttt ttccttggga aaagtgtccg 780
218 acagaaccaa ggctcggtaa aggatactaa taatgtaata atataatatt aacaaacatc 840
219 tactgagett taaatacgtg geaggeactg tgetgtgeac tttatgtgea ttateeaaat 900
220 taactctcag taattccagg agttcttatc accggggtct gtggaagagg aaatggtggt 960
221 ttacggcaat ttccgaatgt cgcacagtag aaagtggacc agcaaccagg gttgcagatc 1020
222 ccaaaqtccq qqatcttcac catttcaccc aaccgcctta gagacctccc agcaaaaggt 1080
223 ctaagcacga atgctccaga aaacaggtcc aggtcacgcg tgccttctcc ctttccctct 1140
224 gggctcgaca gtggtgcaca gagcgactag ggagtggggg ccgcggttat ttcattagaa 1200
225 aagetetgga geeeegggag ceettegtee eaggegeact tgeeetgeaa geeettetgg 1260
226 aaggeggeag egtecaggeg gtecagegea tegageegeg ceaggegeac egagatgeec 1320
227 ccgaatctgt ccagctcgcc ctgcagatcg cacgctccaa ctggcggatt ccgcacgtaa 1380
228 ccctgtgcgc ccgaggccca gcggtaaccc gccgaaggcc cggggccgta ggttcgggca 1440
229 agcategege gecagegee ceageteage ggetgeegea tetecaegee gettaatteg 1500
230 teegttgeec aaatgaceee teegegetee agageggaga tegeggeegt acattttaga 1560
231 agagtgccat caataccctc agagttgagc tggggttacc actctgcaga gctatgccaa 1620
232 ggagaccgcg aagtggtcca ccagcccttc tcagcaattt gccccgctcc agttaagcgg 1680
233 ggaaaacctc ctcccccacc gccactcact tattttgttt ggtagtttaa ttcaacagac 1740
234 qttqqtttaq qqatqqqqaa gccgqqaqqa gatqcaaaaa cgtcctgttt ttcaggaatt 1800
235 tacaatctaa qtqqcaqttc cqttcctqqa acctatcttc tttcacaqat gccgttacag 1860
236 ctgtttcaaa gactgctggt tttgaaaacg ctcagtccct cccaccacgc tgtgcactgc 1920
237 tgcgctccac tcaggggatt acggcgcagg ggcggaccct cgctgacttc tgccccggaa 1980
238 gtttttctct cagttgaagc gcgcacattg agtcggcttt tctactgctt cggctagggt 2040
239 accttgtgac catgtcttcc aagaagaata gaaagcggtt gaaccaaagc gcggaaaatg 2100
240 gttcgtcctt gccctctgct gcttcctctt gtgcggaggc acgggctcct tctgctggat 2160
241 cagacttege ggeaacetee gggaetetga eggtgaecaa ettattagaa aagggtaaag 2220
242 aattccqqqt qqqaqactaq aggccqaggg ggcgcaacct gaggggcggg agactcagtg 2280
243 atactgactt ggggtgcaga gacttttgaa agttgggcgt ggcatgggtc tgagagcggc 2340
244 acagaagegt gtaagactte tgtatgtaag aageatgeaa aegtageege geetgeactg 2400
245 gagaagttag gctgaaatgg tttatccgtg tccaaggttt ctcttttgta aattattttg 2460
246 aatttttaag ttgcacttga aacgctttgt ctcatttaac attaacgaca catttaattg 2520
247 caaaattgtt cataactagc gaacatggga tacttatcct aaaaaactaa ataaccaatt 2580
248 cagtaaaaat gtggatcact ctgcctttag gagtgatgat agcattttct tccagtacta 2640
249 cqcaqqttat qatatttaat aactqctaqt aactqcacac tagtqttagc agtatatata 2700
250 gctcatctaa tttctacaga agccgtgacc gatgggtata gtaatgacag ttttatagag 2760
251 ggaggaaagt taacgtgatt ccattcaggc tacacaggtg ctaagtggca gagccgatat 2820
252 ttgagtccag atgtgtttga cttaaaagcc attcctagat gacagctcct ttgatttggg 2880
253 agaacgtatt tatactgaaa agtcttataa attgaattaa aaattgaaaa tttatatttg 2940.
254 atgtgtaata gctttgaaaa atggaccaga ctagtggcaa aaacttttat gtctaatcag 3000
255 accttaaaga caatttagag ttcacatggt attttcactc tgccttaggt atcagaggac 3060
256 aacttettet etecteaagg ticaacttet tittegaacta gateetteee attegacatet 3120 · .
```

Input Set : A:\SEQLIST_1361US.txt

Output Set: N:\CRF4\09092004\J506549.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

```
Seq#:3; N Pos. 10891,10892,10893,10894,10895,10896,10897,10898,10899,10900
Seq#:3; N Pos. 10901,10902,10903,10904,10905,10906,10907,10908,10909,10910
Seq#:3; N Pos. 10911,10912,10913,10914,10915,10916,10917,10918,10919,10920
Seq#:3; N Pos. 10921,10922,10923,10924,10925,10926,10927,10928,10929,10930
Seq#:3; N Pos. 10931,10932,10933,10934,10935,10936,10937,10938,10939,10940
Seq#:3; N Pos. 10941,10942,10943,10944,10945,10946,10947,10948,10949,10950
Seq#:3; N Pos. 10951,10952,10953,10954,10955,10956,10957,10958,10959,10960
Seq#:3; N Pos. 10961,10962,10963,10964,10965,10966,10967,10968,10969,10970
Seq#:3; N Pos. 10971,10972,10973,10974,10975,10976,10977,10978,10979,10980
Seq#:3; N Pos. 10981,10982,10983,10984,10985,10986,10987,10988,10989,10990
Seq#:3; N Pos. 10991,10992,10993,10994,10995,10996,10997,10998,10999,11000
Seq#:3; N Pos. 11001,11002,11003,11004,11005,11006,11007,11008,11009,11010
Seq#:3; N Pos. 11011,11012,11013,11014,11015,11016,11017,11018,11019,11020
Seq#:3; N Pos. 11021,11022,11023,11024,11025,11026,11027,11028,11029,11030
Seq#:3; N Pos. 11031,11032,11033,11034,11035,11036,11037,11038,11039,11040
Seq#:3; N Pos. 11041,11042,11043,11044,11045,11046,11047,11048,11049,11050
Seq#:3; N Pos. 11051,11052,11053,11054,11055,11056,11057,11058,11059,11060
Seq#:3; N Pos. 11061,11062,11063,11064,11065,11066,11067,11068,11069,11070
Seq#:3; N Pos. 11071,11072,11073,11074,11075,11076,11077,11078,11079,11080
Seq#:3; N Pos. 11081,11082,11083,11084,11085,11086,11087,11088,11089,11090
Seq#:3; N Pos. 11091,11092,11093,11094,11095,11096,11097,11098,11099,11100
Seq#:3; N Pos. 11101,11102,11103,11104,11105,11106,11107,11108,11109,11110
Seq#:3; N Pos. 11111,11112,11113,11114,11115,11116,11117,11118
Seq#:3; N Pos. 67140,67141,67142,67143,67144,67145,67146,67147,67148,67149
Seq#:3; N Pos. 67150,67151,67152,67153,67154,67155,67156,67157,67158,67159
Seq#:3; N Pos. 77171,77172,77173,77174,77175,77176,77177,77178,77179,77180
Seq#:3; N Pos. 77181,77182,77183,77184,77185,77186,77187,77188,77189,77190
Seq#:3; N Pos. 77191,77192,77193,77194,77195,77196,77197,77198,77199,77200
Seq#:3; N Pos. 77201,77202,77203,77204,77205,77206,77207,77208,77209,77210
Seq#:3; N Pos. 77211,77212,77213,77214,77215,77216,77217,77218,77219,77220
Seq#:3; N Pos. 77221,77222,77223,77224,77225,77226,77227,77228,77229,77230
Seq#:3; N Pos: 77231,77232,77233,77234,77235,77236,77237,77238,77239,77240
Seq#:3; N Pos. 77241,77242,77243,77244,77245,77246,77247,77248,77249,77250
Seq#:3; N Pos. 77251,77252,77253,77254,77255,77256,77257,77258,77259,77260
Seq#:3; N Pos. 77261,77262,77263,77264,77265,77266,77267,77268,77269,77270
Seq#:3; N Pos. 77271,77272,77273,77274,77275,77276,77277,77278,77279,77280
Seq#:3; N Pos. 77281,77282,77283,77284,77285,77286,77287,77288,77289,77290
Seq#:3; N Pos. 77291,77292,77293,77294,77295,77296,77297,77298,77299,77300
Seq#:3; N Pos. 77301,77302,77303,77304,77305,77306,77307,77308,77309,77310
Seq#:3; N Pos. 77311,77312,77313,77314,77315,77316,77317,77318,77319,77320
Seq#:3; N Pos. 77321,77322,77323,77324,77325,77326,77327,77328,77329,77330
Seq#:3; N Pos. 77331,77332,77333,77334,77335,77336,77337,77338,77339,77340
Seq#:3; N Pos. 77341,77342,77343,77344,77345,77346,77347,77348,77349,77350
Seq#:3; N Pos. 77351,77352,77353,77354,77355,77356,77357,77358,77359,77360
```

Input Set : A:\SEQLIST_1361US.txt

Output Set: N:\CRF4\09092004\J506549.raw

Seq#:3; N Pos. 77361,77362,77363,77364,77365,77366,77367,77368,77369,77370
Seq#:3; N Pos. 77371,77372,77373,77374,77375,77376,77377,77378,77379,77380
Seq#:3; N Pos. 77381,77382,77383,77384,77385,77386,77387,77388,77389,77390
Seq#:3; N Pos. 77391,77392,77393,77394,77395,77396,77397,77398,77399,77400
Seq#:3; N Pos. 77401,77402,77403,77404,77405,77406,77407,77408,77409,77410
Seq#:3; N Pos. 77411,77412,77413,77414,77415,77416,77417,77418,77419,77420
Seq#:3; N Pos. 77421,77422,77423,77424,77425,77426,77427,77428,77429,77430

VERIFICATION SUMMARY

DATE: 09/09/2004

PATENT APPLICATION: US/10/506,549

TIME: 16:15:51

Input Set : A:\SEQLIST 1361US.txt

Output Set: N:\CRF4\09092004\J506549.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application Number

L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:386 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:10860

M:341 Repeated in SeqNo=3